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RAW SEQUENCE LISTING PATENT APPLICATION US/09/016,743

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This Raw Listing contains the General Information Section and up to the first 5 pages.

SEQUENCE LISTING ENTERED (1) General Information: (i) APPLICANT: Rosenblatt, Joseph D. Challita-Eid, Pia 6 7 Morrison, Sherie Abboud, Camille N. 8 9 Shin, Seung-Uon 10 (ii) TITLE OF INVENTION: CHIMERIC ANTIBODY FUSION PROTEINS FOR 11 THE RECRUITMENT AND STIMULATION OF AN ANTITUMOR IMMUNE 12 13 RESPONSE 14 (iii) NUMBER OF SEQUENCES: 10 15 16 (iv) CORRESPONDENCE ADDRESS: 17 (A) ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP 18 (B) STREET: Clinton Square, P.O. Box 1051 19 20 (C) CITY: Rochester 21 (D) STATE: New York 22 (E) COUNTRY: USA 23 (F) ZIP: 14603 24 25 (V) COMPUTER READABLE FORM: 26 (A) MEDIUM TYPE: Floppy disk 27 (B) COMPUTER: IBM PC compatible 28 (C) OPERATING SYSTEM: PC-DOS/MS-DOS 29 (D) SOFTWARE: Patentin Release #1.0, Version #1.30 30 (vi) CURRENT APPLICATION DATA: 31 32 (A) APPLICATION NUMBER: 33 (B) FILING DATE: 34 (C) CLASSIFICATION: 35 (vii) PRIOR APPLICATION DATA: 36 (A) APPLICATION NUMBER: US 60/307,256 37 38 (B) FILING DATE: 31-JAN-1997 39 (vii) PRIOR APPLICATION DATA: 40 41 (A) APPLICATION NUMBER: US 60/064,018 42 (B) FILING DATE: 03-NOV-1997 43 (viii) ATTORNEY/AGENT INFORMATION: 44 45 (A) NAME: Goldman, Michael L.

(B) REGISTRATION NUMBER: 30,727





RAW SEQUENCE LISTING PATENT APPLICATION US/09/016,743

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| 47 | (C) REFERENCE/DOCKET NUMBER: 176/60192 |
|-----|--|
| 48 | |
| 49 | (ix) TELECOMMUNICATION INFORMATION: |
| 50 | (A) TELEPHONE: 716-263-1304 |
| 51 | (B) TELEFAX: 716-263-1600 |
| 52 | |
| 53 | |
| 54 | (2) INFORMATION FOR SEQ ID NO:1: |
| 55 | |
| 56 | (i) SEQUENCE CHARACTERISTICS: |
| 57 | (A) LENGTH: 30 base pairs |
| 5.8 | (B) TYPE: nucleic acid |
| 59 | (C) STRANDEDNESS: single |
| 60 | (D) TOPOLOGY: unknown |
| 61 | |
| 62 | (ii) MOLECULE TYPE: other nucleic acid |
| 63 | (A) DESCRIPTION: /desc = "DNA Primer" |
| 64 | And the second of the second o |
| 65 | |
| 66 | |
| 67 | |
| 68 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1: |
| 69 | (XI) SEQUENCE DESCRIPTION. SEQ ID NO.1. |
| | GGCATAAGCT TGATATCTGA AGCCATGGGC 30 |
| 70 | GGCATAAGCT TGATATCTGA AGCCATGGGC 30 |
| 71 | (A) TANDONALITON FOR GEO. TO NO. 3. |
| 72 | (2) INFORMATION FOR SEQ ID NO:2: |
| 73 | ALL GROUPING GUARAGERTAGE |
| 74 | (i) SEQUENCE CHARACTERISTICS: |
| 75 | (A) LENGTH: 28 base pairs |
| 76 | (B) TYPE: nucleic acid |
| 77 | (C) STRANDEDNESS: single |
| 78 | (D) TOPOLOGY: unknown |
| 79 | |
| 80 | (ii) MOLECULE TYPE: other nucleic acid |
| 81 | (A) DESCRIPTION: /desc = "DNA Primer" |
| 82 | |
| 83 | |
| 84 | |
| 85 | |
| 86 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: |
| 87 | |
| 88 | GCGCGGTTAA CCGTTATCAG GAAAATGC 28 |
| 89. | |
| 90 | (2) INFORMATION FOR SEQ ID NO:3: |
| 91 | |
| 92 | (i) SEQUENCE CHARACTERISTICS: |
| 93 | (A) LENGTH: 39 base pairs |
| 94 | (B) TYPE: nucleic acid |
| 95 | (C) STRANDEDNESS: single |
| 96 | (D) TOPOLOGY: unknown |
| 97 | |
| 98 | (ii) MOLECULE TYPE: other nucleic acid |
| 99 | (A) DESCRIPTION: /desc = "DNA Primer" |

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| 100 | |
|-----|--|
| 101 | |
| 102 | |
| 103 | |
| 104 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: |
| 105 | |
| 106 | GGGGATATCC ACCATGGRAT GSAGCTGKGT MATSCTCTT 3 |
| 107 | |
| 108 | (2) INFORMATION FOR SEQ ID NO:4: |
| 109 | · · · · · · · · · · · · · · · · · · · |
| 110 | (i) SEQUENCE CHARACTERISTICS: |
| 111 | (A) LENGTH: 39 base pairs |
| 112 | (B) TYPE: nucleic acid |
| 113 | (C) STRANDEDNESS: single |
| 114 | (D) TOPOLOGY: unknown |
| | (D) TOPOLOGI: unknown |
| 115 | And the second makes the second secon |
| 116 | (ii) MOLECULE TYPE: other nucleic acid |
| 117 | (A) DESCRIPTION: /desc = "DNA Primer" |
| 118 | |
| 119 | |
| 120 | |
| 121 | |
| 122 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: |
| 123 | |
| 124 | GACTGGGTCA TCTGGATGTC GGAGTGGACA CCTGTGGAG |
| 125 | |
| 126 | (2) INFORMATION FOR SEQ ID NO:5: |
| 127 | |
| 128 | (i) SEQUENCE CHARACTERISTICS: |
| 129 | (A) LENGTH: 38 base pairs |
| 130 | (B) TYPE: nucleic acid |
| 131 | (C) STRANDEDNESS: single |
| 132 | (D) TOPOLOGY: unknown |
| 133 | (b) Iotoboti dilinomi |
| 134 | (ii) MOLECULE TYPE: other nucleic acid |
| 135 | (A) DESCRIPTION: /desc = "DNA Primer" |
| 136 | (A) DESCRIPTION: / desc - DNA FILMET |
| | |
| 137 | |
| 138 | |
| 139 | |
| 140 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: |
| 141 | |
| 142 | CTCCACAGGT GTCCACTCCG ACATCCAGAT GACCCAGT |
| 143 | |
| 144 | (2) INFORMATION FOR SEQ ID NO:6: |
| 145 | |
| 146 | (i) SEQUENCE CHARACTERISTICS: |
| 147 | (A) LENGTH: 32 base pairs |
| 148 | (B) TYPE: nucleic acid |
| 149 | (C) STRANDEDNESS: single |
| 150 | (D) TOPOLOGY: unknown |
| 151 | |
| 152 | (ii) MOI FOULE TYPE: other nucleic acid |

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| 153 | (A) DESCRIPTION: /desc = "DNA Primer" | |
|-------|--|-------------------|
| 154 | | • |
| 155 | | • |
| 156 | | |
| 157 | | |
| 158 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6: | - W |
| 159 | | · · · · · · · · · |
| 160 | GCTTGTCGAC TTACGTTTGA TCTCCACCTT GG | 32 |
| 161 | | |
| 162 | (2) INFORMATION FOR SEQ ID NO:7: | |
| 163 | | |
| 164 | (i) SEQUENCE CHARACTERISTICS: | |
| : 165 | (A) LENGTH: 39 base pairs | |
| 166 | (B) TYPE: nucleic acid | |
| 167 | (C) STRANDEDNESS: single | * |
| 168 | (D) TOPOLOGY: unknown | |
| 169 | | |
| 170 | (ii) MOLECULE TYPE: other nucleic acid | |
| 171 | (A) DESCRIPTION: /desc = "DNA Primer" | |
| 172 | | 5 |
| 173 | | • • |
| 174 | | |
| 175 | | |
| 176 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7: | |
| 177 | | |
| 178 | GACTCCACCA GCTGAACCTC GGAGTGGACA CCTGTGGAG | 39 |
| 179 | | , |
| 180 | (2) INFORMATION FOR SEQ ID NO:8: | |
| 181 | | |
| 182 | (i) SEQUENCE CHARACTERISTICS: | |
| 183 | (A) LENGTH: 38 base pairs | |
| 184 | (B) TYPE: nucleic acid | |
| 185 | (C) STRANDEDNESS: single | • |
| 186 | (D) TOPOLOGY: unknown | |
| 187. | | ., |
| 188 | (ii) MOLECULE TYPE: other nucleic acid | |
| 189 | (A) DESCRIPTION: /desc = "DNA Primer" | |
| 190 | | |
| 191 | | * . |
| 192 | | |
| 193 | | |
| 194 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8: | |
| 195 | | |
| 196 | CTCCACAGGT GTCCACTCCG AGGTTCAGCT GGTGGAGT | 38 |
| 197 | | |
| 198 | (2) INFORMATION FOR SEQ ID NO:9: | |
| 199 | | |
| 200 | (i) SEQUENCE CHARACTERISTICS: | 20 |
| 201 | (A) LENGTH: 29 base pairs | |
| 202 | (B) TYPE: nucleic acid | |
| 203 | (C) STRANDEDNESS: single | Þ |
| 204 | (D) TOPOLOGY: unknown | |
| 205 | | *. |

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| 206 | (ii) MOLECULE TYPE: other nucleic acid | |
|------------|--|-----------------------|
| 207 | (A) DESCRIPTION: /desc = "DNA Primer" | • |
| 208 209 | | |
| 210 | | |
| 211 | | |
| 212 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: | |
| 213 | (11) 52252102 5250111 12011 122 12 10013 | |
| 214 | TTGGTGCTAG CCGAGGAGAC GGTGACCAG | 2: |
| 215 | | |
| 216 | (2) INFORMATION FOR SEQ ID NO:10: | |
| 217 | | * |
| 218 | (i) SEQUENCE CHARACTERISTICS: | * |
| 219 | | 40 0 |
| 220 | (B) TYPE: nucleic acid | * |
| 221 | (C) STRANDEDNESS: single | |
| 222 | (D) TOPOLOGY: unknown | |
| 223 | | and the second second |
| 224 | (ii) MOLECULE TYPE: other nucleic acid | • |
| 225 226 | (A) DESCRIPTION: /desc = "DNA Primer" | |
| 226 | | • |
| 227 | | = |
| 229 | | * |
| 230 | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: | |
| 231 | // | |
| 232 | GGCGGCGGAT CCGAGGTTCA GCTGGTG | 2 |
| 000 | | |







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